MLLNVLRICI	IVCLVNDGAG	KHSEGRERTK	TYSLNSRGYF	40
RKERGARRSK	ILLVNTKGLD	EPHIGHGDFG	LVAELFDSTR	80
THTNRKEPDM	NKVKLFSTVA	HG <u>NKS</u> ARRKA	YNGSRRNIFS	120
RRSFDKRNTE	VTEKPGAKMF	WNNFLVKMNG	APQ <u>NTS</u> HGSK	160
AQEIMKEACK	TLPFTQNIVH	ENCDRMVIQN	NLCFGKCISL	200
HVPNQQDRRN	TCSHCLPSKF	TLNHLTLNCT	GSKNVVKVVM	240
MVEECTCEAH	KSNFHQTAQF	NMDTSTTLHH		270

Figure 1. Deduced amino acid sequence of Xenopus cerberus protein. SEQ ID NO:1.

Figure 2. Nucleotide sequence of the full-length cerberus DNA derived from the Xenopus organizer. The sense strand is on top (in the 5' to 3' direction) and the antisense strand on the bottom line (on the opposite direction). SEQ ID NO:2.

GAATTCCCAG CAAGTCGCTC AGAAACACTG CAGGGTCTAG ATATCATACA ATGTTACTAA	60
CTTAAGGGTC GTTCAGCGAG TCTTTGTGAC GTCCCAGATC TATAGTATGT TACAATGATT	
ATGTACTCAG GATCTGTATT ATCGTCTGCC TTGTGAATGA TGGAGCAGGA AAACACTCAG	120
TACATGAGTC CTAGACATAA TAGCAGACGG AACACTTACT ACCTCGTCCT TTTGTGAGTC	
AAGGACGAGA AAGGACAAAA ACATATTCAC TTAACAGCAG AGGTTACTTC AGAAAAGAAA	180
TICCIGCTCT TICCIGITIT TGTATAAGTG AATTGTCGTC TCCAATGAAG TCTTTTCTTT	
TICCIGCICI TICCIGITII IGIRIRAGIO MATICIOGIO TOMINICALI	
GAGGAGCACG TAGGAGCAAG ATTCTGCTGG TGAATACTAA AGGTCTTGAT GAACCCCACA	240
CTCCTCGTGC ATCCTCGTTC TAAGACGACC ACTTATGATT TCCAGAACTA CTTGGGGTGT	
TTGGGCATGG TGATTTTCGC TTAGTAGCTG AACTATTTGA TTCCACCAGA ACACATACAA	300
AACCCGTACC ACTAAAAGCG AATCATCGAC TTGATAAACT AAGGTGGTCT TGTGTATGTT	
ACAGAAAAGA GCCAGACATG AACAAAGTCA AGCTTTTCTC AACAGTTGCC CATGGAAACA	360
TGTCTTTCT CGGTCTGTAC TTGTTTCAGT TCGAAAAGAG TTGTCAACGG GTACCTTTGT	
TGTCTTTTCT CGGTCTGTAC TTGTTTCAGT TCGAAAAGAG TTGTCGGGGGGG	
AAAGTGCAAG AAGAAAAGCT TACAATGGTT CTAGAAGGAA TATTTTTCCT CGCCGTTCTT	420
TTTCACGTTC TTCTTTTCGA ATGTTACCAA GATCTTCCTT ATAAAAAGGA GCGGCAAGAA	
TTGATAAAAG AAATACAGAG GTTACTGAAA AGCCTGGTGC CAAGATGTTC TGGAACAATT	480
AACTATTTC TTTATGTCTC CAATGACTTT TCGGACCACG GTTCTACAAG ACCTTGTTAA	
	540
TTTTGGTTAA AATGAATGGA GCCCCACAGA ATACAAGCCA TGGCAGTAAA GCACAGGAAA AAAACCAATT TTACTTACCT CGGGGTGTCT TATGTTCGGT ACCGTCATTT CGTGTCCTTT	340
AAAACCARTT TTACTIACCI CGGGGIGICI IRIGIICGGI ACCGICATTI GGIGGGG	
TAATGAAAGA AGCTTGCAAA ACCTTGTTTT TCACTCAGAA TATTGTACAT GAAAACTGTG	600
ATTACTTTCT TCGAACGTTT TGGAACAAAA AGTGAGTCTT ATAACATGTA CTTTTGACAC	
ACAGGATGGT GATACAGAAC AATCTGTGCT TTGGTAAATG CATCTCTCTC CATGTTCCAA	660
TGTCCTACCA CTATGTCTTG TTAGACACGA AACCATTTAC GTAGAGAGAG GTACAAGGTT	
	720
ATCAGCAAGA TCGACGAAAT ACTTGTTCCC ATTGCTTGCC GTCCAAATTT ACCCTGAACC TAGTCGTTCT AGCTGCTTTA TGAACAAGGG TAACGAACGG CAGGTTTAAA TGGGACTTGG	120
TAGTOGTTOT AGOTGOTTA TGAACAAGGG TAACGAACGG CAGGTTTAAA 1000001100	
ACCTGACGCT GAATTGTACT GGATCTAAGA ATGTAGTAAA GGTTGTCATG ATGGTAGAGG	780
TGGACTGCGA CTTAACATGA CCTAGATTCT TACATCATTT CCAACAGTAC TACCATCTCC	
AATGCACGTG TGAAGCTCAT AAGAGCAACT TCCACCAAAC TGCACAGTTT AACATGGATA	840
TTACGTGCAC ACTTCGAGTA TTCTCGTTGA AGGTGGTTTG ACGTGTCAAA TTGTACCTAT	
CATCTACTAC CCTGCACCAT TAAAGGACTG CCATACAGTA TGGAAATGCC CTTTTGTTGG	900
CATCTACTAC CCTGCACCAT TAAAGGACTG CCATACAGIA IGGARATGCC CTITIOTICG GTAGATGATG GGACGTGGTA ATTTCCTGAC GGTATGTCAT ACCTTTACGG GAAAACAACC	,
GINGNIGNIG GGNCGIGGIN MILLOCIGNO GGINIGIONI NOOTTINGGO GERNANDA	
ARTATTTGTT ACATACTATG CATCTAAAGC ATTATGTTGC CTTCTATTTC ATATAACCAC	960
TTATAAACAA TGTATGATAC GTAGATTTCG TAATACAACG GAAGATAAAG TATATTGGTG	í
•	
ATGGAATAAG GATTGTATGA ATTATAATTA ACAAATGGCA TTTTGTGTAA CATGCAAGAT	1020
TACCTTATTC CTAACATACT TAATATTAAT TGTTTACCGT AAAACACATT GTACGTTCTA	<b>L</b>

CTCTGTTCCA GAGACAAGGT	TCAGTTGCAA AGTCAACGTT	GATAAAAGGC CTATTTTCCG	AATATTTGTT TTATAAACAA	TGACTTTTT ACTGAAAAAA	TCTACAAAAT AGATGTTTTA	1080
	ATATATGATA TATATACTAT					1140
	TTTGCCCAGG AAACGGGTCC					1200
	TTTAAAAGCA AAATTTTCGT					1260
	TCATAGGGGG AGTATCCCCC					1320
TGTTACAAAA ACAATGTTTT						

Fig. 2. (Continuation page 2, SEQ ID NO:2).

1	MSRTRKVDSL	LLLAIPGLAL	LLLPNAYCAS	CEPVRIPMCK	SMPWNMTKMP	NHLHHSTQAN	60
1	AILAIEQFEG	LLTTECSQDL	LFFLCAMYAP	ICTIDFQHEP	IKPCKSVCER	ARAGCEPILI	120
1	KYRHTWPESL	ACEELPVYDR	GVCISPEAIV	TVEQGTDSMP	DFSMDSNNGN	CGSGREHCKC	180
1	KPMKATQKTY	LKNNYNYVIR	AKVKEVKVKC	HDATAIVEVK	EILKSSLVNI	PKDTVTLYTN	240
į	sgclcpqlva	NEEYIIMGYE	DKERTRLLLV	EGSLAEKWRD	RLAKKVKRWD	QKLRRPRKSK	300
	DPVAPIPNKN	SNSRQARS					

Figure 3. Deduced amino acid sequence of Xenopus frazzled protein. SEQ ID NO:3.

Figure 4. Nucleotide sequence of the full-length frazzled cDNA derived from the Xenopus organizer. The sense strand of the DNA on top (5' to 3' direction) and the antisense strand on the bottom line (opposite direction). SEQ ID NO:4.

GAATTCCCTT	TCACACAGGA	CTCCTGGCAG	aggtgaatgg	TTAGCCCTAT	GGATTTGGTT	60
CTTAAGGGAA	AGTGTGTCCT	GAGGACCGTC '	TCCACTTACC .	AATCGGGATA	CCTAAACCAA	
TGTTGATTTT	GACACATGAT	TGATTGCTTT	CAGATAGGAT	TGAAGGACTT	GGATTTTAT	120
ACAACTAAAA	CTGTGTACTA	ACTAACGAAA	GTCTATCCTA	ACTTCCTGAA	CCTAAAAATA	
CTAATTCTGC	ACTTTTAAAT	TATCTGAGTA	ATTGTTCATT	TTGTATTGGA	TGGGACTAAA	180
GATTAAGACG	TGAAAATTTA	ATAGACTCAT	TAACAAGTAA	AACATAACCT	ACCCTGATTT	
GATAAACTTA	ACTCCTTGCT	TTTGACTTGC	CCATAAACTA	TAAGGTGGGG	TGAGTTGTAG	240
CTATTTGAAT	TGAGGAACGA	AAACTGAACG	GGTATTTGAT	ATTCCACCCC	ACTCAACATC	
TTGCTTTTAC	ATGTGCCCAG	ATTTTCCCTG	TATTCCCTGT	ATTCCCTCTA	AAGTAAGCCT	300
AACGAAAATG	TACACGGGTC	TAAAAGGGAC	ATAAGGGACA	TAAGGGAGAT	TTCATTCGGA	
ACACATACAG	GTTGGGCAGA	ATAACAATGT	CTCGAACAAG	GAAAGTGGAC	TCATTACTGC	360
TGTGTATGTC	CAACCCGTCT	TATTGTTACA	GAGCTTGTTC	CTTTCACCTG	AGTAATGACG	
TACTGGCCAT	ACCTGGACTG	GCGCTTCTCT	TATTACCCAA	TGCTTACTGT	GCTTCGTGTG	420
ATGACCGGTA	TGGACCTGAC	CGCGAAGAGA	ATAATGGGTT	ACGAATGACA	CGAAGCACAC	
AGCCTGTGCG	GATCCCCATG	TGCAAATCTA	TGCCATGGAA	CATGACCAAG	ATGCCCAACC	480
TCGGACACGC	CTAGGGGTAC	ACGTTTAGAT	ACGGTACCTT	GTACTGGTTC	TACGGGTTGG	
ATCTCCACCA	CAGCACTCAA	GCCAATGCCA	TCCTGGCAAT	TGAACAGTTT	GAAGGTTTGC	540
TAGAGGTGGT	GTCGTGAGTT	CGGTTACGGT	AGGACCGTTA	ACTTGTCAAA	CTTCCAAACG	
TGACCACTGA	ATGTAGCCAG	GACCTTTTGT	TCTTTCTGTG	TGCCATGTAT	GCCCCCATTT	600
ACTGGTGACT	TACATCGGTC	CTGGAAAACA	AGAAAGACAC	ACGGTACATA	CGGGGGTAAA	
GTACCATCG	TTTCCAGCAT	GAACCAATTA	AGCCTTGCAA	GTCCGTGTGC	GAAAGGCCA	660
CATGGTAGCT	AAAGGTCGTA	CTTGGTTAAT	TCGGAACGTT	CAGGCACACG	CTTTCCCGGT	
GGGCCGGCTG	G TGAGCCCATT	CTCATAAAGT	ACCGGCACAC	TTGGCCAGAG	AGCCTGGCAT	720
CCCGGCCGA	ACTCGGGTAA	GAGTATTTCA	TGGCCGTGTG	AACCGGTCTC	TCGGACCGTA	
GTGAAGAGC	r GCCCGTATAI	GACAGAGGAG	TCTGCATCTC	CCCAGAGGCT	ATCGTCACAG	780
CACTTCTCG	A CGGGCATATA	CTGTCTCCTC	AGACGTAGAG	GGGTCTCCG!	TAGCAGTGTC	
TGGAACAAG	G AACAGATTC	ATGCCAGACT	TCTCCATGGA	TTCAAACAA	GGAAATTGCG	840
ACCTTGTTC	C TTGTCTAAG1	TACGGTCTGA	AGAGGTACCI	AAGTTTGTT!	A CCTTTAACGC	
GAAGCGGCA	G GGAGCACTG	r aaatgcaago	CCATGAAGG	AACCCAAAA	ACGTATCTCA	900
CTTCGCCGT	C CCTCGTGAC	A TTTACGTTC	GGTACTTCC	3 TTGGGTTTT	C TGCATAGAGT	
AGAATAATT	A CAATTATGT	A ATCAGAGCA	AAGTGAAAG	A GGTGAAAGT	G AAATGCCACG	960
TCTTATTAA	T GTTAATACA	T TAGTCTCGT	TTCACTTTC	T CCACTTTCA	C TTTACGGTGC	
ACGCAACAG	C AATTGTGGA	A GTAAAGGAG	A TTCTCAAGT	C TTCCCTAGT	G AACATTCCTA	1020
TGCGTTGTC	G TTAACACCT	T CATTTCCTC	r aagagttca	G AAGGGATCA	C TTGTAAGGAT	

AAGACACAGT	GACACTGTAC	ACCAACTCAG	GCTGCTTGTG	CCCCAGCTT	GTTGCCAATG	1080
TTCTGTGTCA	CTGTGACATG	TGGTTGAGTC	CGACGAACAC	GGGGGTCGAA	CAACGGTTAC	
	AATTATGGGC					1140
TCCTTATGTA	TTAATACCCG	ATACTTCTGT	TTCTCGCATG	GTCCGAAGAT	GATCACCTTC	
		•				
	CGAAAAATGG					1200
CTAGGAACCG	GCTTTTTACC	TCTCTAGCAG	AACGATTCTT	TCAGTTCGCG	ACCCTAGTTT	
	TCCCAGGAAA					1260
TCGAAGCTGC	AGGGTCCTTT	TCGTTTCTGG	GGCACCGAGG	TTAAGGGTTG	TTTTTGTCGT	
	AGCGCGTAGT					1320
TAAGGTCTGT	TCGCGCATCA	ATCTGATTGC	CTTTCCACAT	ACCTTTGAGA	TACCTGAAAC	
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	TTGCATTGTT			•		1360
TTTGATTCTA	AACGTAACAA	CCTTCTCGTT	TTTTCTTTAA	CGTGATGTCG	TGCAATATAA	
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	GATGTTCTTC					1440
GATAACAAAT	GAIGIICIIC	GACCAAATCA	ACIAACAICA	DEMANDUNDA	MAGAMAMA	
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СУСТСУСТСУ	ATGTCTCAGC	CTAAAGAAGC	TCAATTCATT	TCTGATCAAC	TAATGGTGAC	1560
	TACAGAGTCG					
014.014.01						
AAGTGTTTGA	TACTTGGGGA	AAGTGAACTA	ATTGCAATGG	TAAATCAGAG	AAAAGTTGAC	1620
•	ATGAACCCCT					
CAATGTTGCT	TTTCCTGTAG	ATGAACAAGT	GAGAGATCAC	ATTTAAATGA	TGATCACTTT	1680
GTTACAACGA	AAAGGACATC	TACTTGTTCA	CTCTCTAGTG	TAAATTTACT	ACTAGTGAAA	
CCATTTAATA	CTTTCAGCAG	TTTTAGTTAG	ATGACATGTA	GGATGCACCT	AAATCTAAAT	1740
GGTAAATTAT	GAAAGTCGTC	AAAATCAATC	TACTGTACAT	CCTACGTGGA	TTTAGATTTA	
					AGGTAAATGC	1800
TAAAATAGTA	TTTACTTCTC	GACCAAATCT	GACATACCAG	TGACAACCCT	TCCATTTACG	
					AAATAAAAA	1860
GATGAAACAG	TTAAGACAAA	ATTTTTAACG	GATTTATTTA	. TAATTCAGGA	TTTATTTTT	
AAAAAAAAA						
TTTTTTTTT	TTTTT					

Fig. 4. (Continuation page 2, SEQ ID NO:4).

MLLLFRAIPM LLLGLMVLOT DCEIAOYYID EEEPPGTVIA VLSQHSIFNT TDIPATNFRL 60 120 MKQFNNSLIG VRESDGQLSI MERIDREQIC RQSLHCNLAL DVVSFSKGHF KLLNVKVEVR DINDHSPHFP SEIMHVEVSE SSSVGTRIPL EIAIDEDVGS NSIQNFQISN NSHFSIDVLT 180 RADGVKYADL VLMRELDREI QPTYIMELLA MDGGVPSLSG TAVVNIRVLD FNDNSPVFER 240 STIAVDLVED APLGYLLLEL HATDDDEGVN GEIVYGFSTL ASQEVRQLFK INSRTGSVTL 300 EGQVDFETKQ TYEFEVQAQD LGPNPLTATC KVTVHILDVN DNTPAITITP LTTVNAGVAY 360 IPETATKENF IALISTTDRA SGSNGQVRCT LYGHEHFKLQ QAYEDSYMIV TTSTLDRENI 420 AAYSLTVVAE DLGFPSLKTK KYYTVKVSDE NDNAPVFSKP QYEASILENN APGSYITTVI 480 ARDSDSDQNG KVNYRLVDAK VMGQSLTTFV SLDADSGVLR AVRSLDYEKL KQLDFEIEAA 540 DNGIPQLSTR VQLNLRIVDQ NDNCPVITNP LLNNGSGEVL LPISAPQNYL VFQLKAEDSD 600 EGHNSQLFYT ILROPSRLFA INKESGEVFL KKQLNSDHSE DLSIVVAVYD LGRPSLSTNA 660 TVKFILTDSF PSNVEVVILQ PSAEEQHQID MSIIFIAVLA GGCALLLLAI FFVACTCKKK 720 AGEFKQVPEQ HGTCNEERLL STPSPQSVSS SLSQSESCQL SINTESENCS VSSNQEQHQQ 780 TGIKHSISVP SYHTSGWHLD NCAMSISGHS HMGHISTKVQ WAKEIVTSMT VTLILVENQK 840 RRALSSQCRH KPVLNTQMNQ QGSDMPITIS ATESTRVQKM GTAHCNMKRA IDCLTL

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Figure 5. Deduced amino acid sequence of the Xenopus PAPC (paraxial protocadherin) protein. It encodes a member of the cadherin family of transmembrane proteins that has dorsalizing activity when constructs are injected into Xenopus embryos. SEQ ID NO:5.

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Figure 6. Nucleotide sequence of the full-length PAPC cDNA derived from the Xenopus organizer. The sense strand of the DNA is shown in the top line (in the 5' to 3' direction), and the bottom line shows the antisense strand (opposite orientation). SEQ ID NO:6.

GAATTCCCAG AGATGAACTC CTTGAGATTG TTTTAAATGA CTGCAGGTCT GGAAGGATTC	60
CTTAAGGGTC TCTACTTGAG GAACTCTAAC AAAATTTACT GACGTCCAGA CCTTCCTAAG	
ACATTGCCAC ACTGTTTCTA GGCATGARAR ARCTGCARGT TTCARCTTTG TTTTTGGTGC	120
TGTAACGGTG TGACAAAGAT CCGTACTTTT TTGACGTTCA AAGTTGAAAC AAAAACCACG	
	100
AACTITGATI CITCAAGAIG CIGCTICTCI TCAGAGCCAI TCCAATGCIG CIGITGGGAC	180
TTGAAACTAA GAAGTTCTAC GACGAAGAGA AGTCTCGGTA AGGTTACGAC GACAACCCTG	
2222 CARCAR CARCON CONTRACTOR CARCAR	240
TGATGGTTTT ACAAACAGAC TGTGAAATTG CCCAGTACTA CATAGATGAA GAAGAACCCC ACTACCAAAA TGTTTGTCTG ACACTTTAAC GGGTCATGAT GTATCTACTT CTTCTTGGGG .	
ACTACCAAAA TGTTTGTCTG ACACTTTAAC GGGTCATGAT GIATOLAGIT GILDINGT	
CTGGCACTGT AATTGCAGTG TTGTCACAAC ACTCCATATT TAACACTACA GATATACCTG	300
GACCGTGACA TTAACGTCAC AACAGTGTTG TGAGGTATAA ATTGTGATGT CTATATGGAC	
GACCGIGACA ITAROGICA INCICIOTO ICONO	
CAACCAATTT CCGTCTAATG AAGCAATTTA ATAATTCCCT TATCGGAGTC CGTGAGAGTG	360
GTTGGTTAAA GGCAGATTAC TTCGTTAAAT TATTAAGGGA ATAGCCTCAG GCACTCTCAC	
ATGGGCAGCT GAGCATCATG GAGAGGATTG ACCGGGAGCA AATCTGCAGG CAGTCCCTTC	420
TACCCGTCGA CTCGTAGTAC CTCTCCTAAC TGGCCCTCGT TTAGACGTCC GTCAGGGAAG	
	480
ACTGCAACCT GGCTTTGGAT GTGGTCAGCT TTTCCAAAGG ACACTTCAAG CTTCTGAACG	450
TGACGTTGGA CCGAAACCTA CACCAGTCGA AAAGGTTTCC TGTGAAGTTC GAAGACTTGC	
CTTTCCCACT GAAATAATGC	540
TGAAAGTGGA GGTGAGAGAC ATTAATGACC ATAGCCCTCA CTTTCCCAGT GAAATAATGC ACTTTCACCT CCACTCTCTG TAATTACTGG TATCGGGAGT GAAAGGGTCA CTTTATTACG	• • • •
ACTITICACCT CCACTCTCTG TAATTACTGG TATCGGGAGT GALLGOOTGT	
ATGTGGAGGT GTCTGAAAGT TCCTCTGTGG GCACCAGGAT TCCTTTAGAA ATTGCAATAG	600
TACACCTCCA CAGACTITCA AGGAGACACC CGTGGTCCTA AGGAAATCTT TAACGTTATC	
TACACCIOCA CAGACIII di ilouisiano di ilouisi	
ATGAAGATGT TGGGTCCAAC TCCATCCAGA ACTTTCAGAT CTCAAATAAT AGCCACTTCA	660
TACTTCTACA ACCCAGGTTG AGGTAGGTCT TGAAAGTCTA GAGTTTATTA TCGGTGAAGT	
GCATTGATGT GCTAACCAGA GCAGATGGGG TGAAATATGC AGATTTAGTC TTAATGAGAG	720
CGTAACTACA CGATTGGTCT CGTCTACCCC ACTTTATACG TCTAAATCAG AATTACTCTC	
	780
AACTGGACAG GGAAATCCAG CCAACATACA TAATGGAGCT ACTAGCAATG GATGGGGGTG	760
TTGACCTGTC CCTTTAGGTC GGTTGTATGT ATTACCTCGA TGATCGTTAC CTACCCCCAC	
ATGATAACA	840
TACCATCACT ATCTGGTACT GCAGTGGTTA ACATCCGAGT CCTGGACTTT AATGATAACA ATGGTAGTGA TAGACCATGA CGTCACCAAT TGTAGGCTCA GGACCTGAAA TTACTATTGT	• • • • • • • • • • • • • • • • • • • •
ATGGTAGTGA TAGACCATGA CGTCACCART TGTAGGCTGA GGTTAG	
GCCCAGTGTT TGAGAGAAGC ACCATTGCTG TGGACCTAGT AGAGGATGCT CCTCTGGGAT	900
CGGGTCACAA ACTCTCTTCG TGGTAACGAC ACCTGGATCA TCTCCTACGA GGAGACCCTA	
ACCITITGIT GGAGTTACAT GCTACTGACG ATGATGAAGG AGTGAATGGA GAAATTGTTT	960
TGGAAAACAA CCTCAATGTA CGATGACTGC TACTACTTCC TCACTTACCT CTTTAACAAA	
	1000
ATGGATTCAG CACTTTGGCA TCTCAAGAGG TACGTCAGCT ATTTAAAATT AACTCCAGAA	1020
TACCTAAGTC GTGAAACCGT AGAGTTCTCC ATGCAGTCGA TAAATTTTAA TTGAGGTCTT	

		CAAGCAGACT GTTCGTCTGA		1080
		 TACTTGTAAA ATGAACATTT		1140
		 TACCCCTCTG ATGGGGAGAC		1200
		 GAACTTTATA CTTGAAATAT		1260
		 CTGTACTCTT GACATGAGAA		1320
	- · · · · · · ·	 GATAGTTACC CTATCAATGG		1380
		TGCAGAAGAC ACGTCTTCTG		1440
		TGATGAGAAT ACTACTCTTA		1500
 		 AAATAATGCT TTTATTACGA		1560
		AAATGGCAAA TTTACCGTTT		1620
		ATTTGTTTCT TAAACAAAGA		1680
		AAAACTTAAA TTTTGAATTT		1740
 		CACTCGCGTT GTGAGCGCAA		1800
		TAATCCTCTT ATTAGGAGAA	_	1860
			TTCCAGCTCA AAGGTCGAGT	1920
			CTGAGAGATC GACTCTCTAG	1980
			AAACAATTAA TTTGTTAATT	2040
			GGAAGACCTT CCTTCTGGAA	2100
			TCTAACGTTG AGATTGCAAC	2160

Fig. 6. (Continuation page 2, SEQ ID NO:6).

	TTTGCAACCA					2220
TTCAGCAATA	AAACGTTGGT	AGACGTCTTC	TCGTCGTGGT	CTAGCTATAC	AGGTAATATA	
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GTACTTGTAA	AAAGAAAGCT	GGTGAATTTA	AGCAGGTACC	TGAACAACAC	GGAACATGCA	2340
CATGAACATT	TTTCTTTCGA	CCACTTAAAT	TCGTCCATGG	ACTTGTTGTG	CCTTGTACGT	
	CCTGTTAAGC					2400
TACTTCTTGC	GGACAATTCG	TGGGGTAGAG	GGGTCAGCCA	GAGAAGAAGA	AACAGAGTCA	
	0011000000					0.4.60
	CCAACTCTCC					2460
GACTCAGTAC	GGTTGAGAGG	TAGTTATGAC	TTAGACTCTT	AACGTCGCAC	AGGAGATTGG	
NACACCACCA	TCAGCAAACA	CCCATAAACC	<b>እ</b> ርጥርር እጥርጥር	<b>サ</b> ロサカ ( ) カ サ ( ) サ	TATCACACAT	2520
	AGTCGTTTGT					2020
CTGGTTGGCA	CCTGGACAAT	TGTGCAATGA	GCATAAGTGG	ACATTCTCAC	ATGGGGCACA	2580
GACCAACCGT	GGACCTGTTA	ACACGTTACT	CGTATTCACC	TGTAAGAGTG	TACCCCGTGT	
TTAGTACAAA	GGTACAGTGG	GCAAAGGAGA	TAGTGACTTC	AATGACAGTG	ACTCTGATAC	2640
AATCATGTTT	CCATGTCACC	CGTTTCCTCT	ATCACTGAAG	TTACTGTCAC	TGAGACTATG	
	TCAGAAAAGA					2700
ATCACCTCTT	AGTCTTTTCT	TCTCGTAACT	CGTCGGTTAC	GTCCGTGTTC	GGTCACGAGT	
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CAAGGGTCCA	GAAAATGGGA	ACTGCACATT	GCAATATGAA	AAGGGCTATA	GACTGTCTTA	2820
GTTCCCAGGT	CTTTTACCCT	TGACGTGTAA	CGTTATACTT	TTCCCGATAT	CTGACAGAAT	
•	CCTGTATATT					2880
GAGACATCGA	GGACATATAA	TGTTATGGAT	GGTACGTTCT	TACGGATTGG	ACGTGTATGG	
	000000000000000000000000000000000000000	Omm1 mm1 OO1	#1# <b>011</b> #11#	COMCMMCOM)	1.00001.00001	2940
	CTTAGAGACC GAATCTCTGG					2940
CITGGIAIGG	GARICICIGG	GAATAATGGT	AIAGITATTA	GGACAACGAI	INGCCINCGI	
GGCGGAATAT	GAAAGAGATT	TAGTCAACAG	AAGTGCAACG	TTATCTCCGC	AGAGATCGTC	3000
	CTTTCTCTAA					
TAGCAGATAC	CAAGAATTCA	ATTACAGTCC	GCAGATATCA	AGACAGCTTC	ATCCTTCAGA	3060
ATCGTCTATG	GTTCTTAAGT	TAATGTCAGG	CGTCTATAGT	TCTGTCGAAG	TAGGAAGTCT	
					GCAAGTGCTT	3120
TTAACGATGT	' TGGAAAATTA	GTAATCCGTA	CGTTCACTCT	TACGTGTTTC	CGTTCACGAA	
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					TAAAAAACAA	
					CTAACTAGCA	3300
AAAAAATGTA	AAAATAAAT	GGACTTAACT	TACACTGTAA	CAGGACAGTG	GATTGATCGT	

Fig. 6. (Continuation page 3, SEQ ID NO:6).

		TGAAACAGCA ACTTTGTCGT	3360
 	 	TGCCCTCTGT ACGGGAGACA	3420
 		GCATCTCACC CGTAGAGTGG	3480
	 	TCTGTGTTGT AGACACAACA	3540
 =		CCATTCAGAT GGTAAGTCTA	 3600
 	 -	TCAATAAATA AGTTATTTAT	

Fig. 6. (Continuation page 4, SEQ ID NO:6).

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MVCCGPGRML LGWAGLLVLA ALCLLQVPGA QAAACEPVRI PLCKSLPWNM TKMPNHLHHS 60

TQANAILAME QFEGLLGTHC SPDLLFFLCA MYAPICTIDF QHEPIKPCKS VCERARQGCE 120

PILIKYRHSW PESLACDELP VYDRGVCISP EAIVTADGAD FPMDSSTGHC RGASSERCKC 180

KPVRATQKTY FRNNYNYVIR AKVKEVKMKC HDVTAVVEVK EILKASLVNI PRDTVNLYTT 240

SGCLCPPLTV NEEYVIMGYE DEERSRLLLV EGSIAEKWKD RLGKKVKRWD MKLRHLGLGK 300

TDASDSTQNQ KSGRNSNPRP ARS.

Figure 7. Deduced amino acid sequence of mouse FRZB-1 protein. SEQ ID NO:7.

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Figure 8. Nucleotide sequence of the full-length mouse FRZB-1 cDNA. SEQ ID NO:8.

AAGCCTGGGA						60
TTCGGACCCT	GGTACCAGAC	GACGCCGGGC	CCTGCCTACG	ACGATCCTAC	CCGGCCCAAC	
CTAGTCCTGG	CTGCTCTCTG	CCTGCTCCAG	GTGCCCGGAG	CTCAGGCTGC	AGCCTGTGAG	120
GATCAGGACC						
CCTGTCCGCA						180
GGACAGGCGT	AGGGCGACAC	GTTCAGGGAA	GGGACCTTGT	ACTGGTTCTA	CGGGTTGGTG	
CTGCACCACA	GCACCCAGGC	TAACGCCATC	CTGGCCATGG	AACAGTTCGA	AGGGCTGCTG	240
GACGTGGTGT	CGTGGGTCCG	ATTGCGGTAG	GACCGGTACC	TTGTCAAGCT	TCCCGACGAC	
GGCACCCACT						300
CCGTGGGTGA	CGTCGGGCCT	AGAAGAGAAG	AAGGAGACAC	GTTACATGCG	TGGGTAAACG	
ACCATCGACT	TCCAGCACGA	GCCCATCAAG	CCCTGCAAGT	СТСТСТСТСА	GCGCGCCCGA	360
			GGGACGTTCA			500
			CGCCACTCGT			420
GTCCCGACGC	TCGGGTAAGA	GTAGTTCATG	GCGGTGAGCA	CCGGCCTTTC	GAACCGGACG	
GACGACCTCC	СССФСФАССА	CCCCCCCTC	TGCATCTCTC	CTC3CCCC3T	<u> </u>	480
			ACGTAGAGAG			400
GACGGAGCGG	ATTTTCCTAT	GGATTCAAGT	ACTGGACACT	GCAGAGGGGC	AAGCAGCGAA	540
CTGCCTCGCC	TAAAAGGATA	CCTAAGTTCA	TGACCTGTGA	CGTCTCCCCG	TTCGTCGCTT	
<b>ሮሮ</b> ሞሞርር እ አ አ ጥ		CACACCTACA	CAGAAGACCT	አመመመር ርርር አ አ	<u>ርአ አምጥአር</u> አእር	600
			GTCTTCTGGA			000
TATGTCATCC	GGGCTAAAGT	TAAAGAGGTA	AAGATGAAAT	GTCATGATGT	GACCGCCGTT	660
ATACAGTAGG	CCCGATTTCA	ATTTCTCCAT	TTCTACTTTA	CAGTACTACA	CTGGCGGCAA	
CMCC3 3 CMC3	3 CC 3 3 3 MMCM	3 3 3 C C C 3 T C 3	CTGGTAAACA	መመርርን አርርርን	ርአርርርጥር አለጥ	720
			GACCATTTGT			120
	••••		CCACTTACTG			780
GAAATATGGT	GGAGACCGAC	GGAGACAGGA	GGTGAATGAC	AGTTACTCCT	TATACAGTAG	
\$ mccccm \$ mc	********	እ <i>ር</i> ርመሙርርእርር	ምጥ እ ርጥር ጥጥርር	ጥልር አልርርርጥር	TATAGCTGAG	840
					ATATCGACTC	040
					CCGACACCTT	900
TTCACCTTCC	TAGCCGAACC	ATTCTTTCAG	TTCGCGACCC	TATACTTTGA	GGCTGTGGAA	
GG I CTCCCTA	<b>አ አ አ</b> ርጥር አጥር ር	' സമദേഗദമ <i>സ</i> സഹ	· ልርጥሮልፎልልጥር	' <b>እ</b> ርል እርጥርጥርር	CAGGAACTCT	960
					GTCCTTGAGA	200

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	CAGCACGCAG GTCGTGCGTC					1020
	GGCGCTGGTG CCGCGACCAC					1080
	CAGACACCGC GTCTGTGGCG					1140
	TGGGGTTAGA ACCCCAATCT					1200
	TTTTGCAACC AAAACGTTGG					1260
	TGGGTTTAAT ACCCAAATTA					1320
					AGCCTCTGCT TCGGAGACGA	1380
					TGTGAAAGTT ACACTTTCAA	1440
					GCAGAGTAGC CGTCTCATCG	1500
					TCAAGATTGG AGTTCTAACC	1560
					GGTTAAGGGG CCAATTCCCC	1620
CACACTGGAA GTGTGACCTT	TCAGTAGCCC AGTCATCGGG	TTGAGCCATT AACTCGGTAA	AACAGCAGTG	TTCTTCTGGC	AAGTTTTTGA TTCAAAAACT	1680
TTTGTTCATA AAACAAGTAT	AATGTATTCA TTACATAAGT	CGAGCATTAG CGCTCGTAATG	AGATGAACTT TCTACTTGAA	TATAACTAGAC	TAGACAACAA	1740
ATCTCTATAC TAGAGATATC	CTCTGCTTCC	TTCTAAATCA AAGATTTAG	A AACCCATTGT TTGGGTAACA	TGGATGCTCC	CTCTCCATTC GAGAGGTAAG	1800

	TTGGCTTGCT AACCGAACGA	 	 	1860
	GTGTTATTTA CACAATAAAT	 	 	1920
ACACGGAAAT TGTGCCTTTA	GTGCACATTT CACGTGTAAA	 TTTCTTCCTT AAAGAAGGAA		1980
	TGTGTTTATG ACACAAATAC	 		2040
	ACTAGATTAG TGATCTAATC	 		2100
	TAATGCTCCA ATTACGAGGT	 	 	2160
CGACAACAAC	AACAAA			

CGACAACAAC AACAAA GCTGTTGTTG TTGTTT

MVCGSPGGML	LLRAGLLALA	ALCLLRVPGA	RAAACEPVRI	PLCKSLPWNM	TKMPNHLHHS	60
TQANAILAIE	QFEGLLGTHC	SPDLLFFLCA	MYAPICTIDF	QHEPIKPCKS	VCERARQGCE	120
PILIKYRHSW	PENLACEELP	VYDRGVCISP	EAIVTADGAD	FPMDSSNGNC	RGASSERCKC	180
KPIRATQKTY	FRNNYNYVIR	AKVKEIKTKC	HDVTAVVEVK	EILKSSLVNI	PRDTVNLYTS	240
SGCLCPPLNV	NEEYIIMGYE	DEERSRLLLV	EGSIAEKWKD	RLGKKVKRWD	MKLRHLGLSK	300
SDSSNSDSTQ	SQKSGRNSNP	RQARN.				

Figure 9. Deduced amino acid sequence of human FRZB-1 protein. SEQ ID NO:9.

Figure 10. Nucleotide sequence of the full-length human FRZB-1 cDNA. SEQ ID NO:10. This sequence was assembled from public ESTs from the Genbank database (accession numbers: H18848, R63748, W38677, W44760, H38379 and N71244).

GGCGGAGCGG G	CCTTTTGGC	GTCCACTGCG	CGGCTGCACC	CTGCCCCATC	TGCCGGGATC	60
CCGCCTCGCC C	CGGAAAACCG	CAGGTGACGC	GCCGACGTGG	GACGGGGTAG	ACGGCCCTAG	
ATGGTCTGCG C	GCAGCCCGGG	AGGGATGCTG	CTGCTGCGGG	CCGGGCTGCT	TGCCCTGGCT	120
TACCAGACGC C	CGTCGGGCCC	TCCCTACGAC	GACGACGCCC	GGCCCGACGA	ACGGGACCGA	
GCTCTCTGCC 7						180
CGAGAGACGG A	ACGAGGCCCA	CGGGCCCCGA	GCCCGACGTC	GGACACTCGG	GCAGGCGTAG	
CCCCTGTGCA A	N CERCOCERCOC	CMCCA A CAMC	A CMA A CAMCC	CCXXCCXCCT	CCACCACACC	240
GGGGACACGT						240
GGGGACACGI I	DEDARDERATI	GACCIIGIAC	IGATICIACG	GGTTGGTGGA	0010010100	
ACTCAGGCCA A	ACGCCATCCT	GGCCATCGAG	CAGTTCGAAG	GTCTGCTGGG	CACCCACTGC	300
TGAGTCCGGT						
IGNOTOCOGI .	rocogracoa	ccoornocre	01011100110	00		
AGCCCCGATC T	TGCTCTTCTT	CCTCTGTGCC	ATGTACGCGC	CCATCTGCAC	CATTGACTTC	360
TCGGGGCTAG A						
CAGCACGAGC (	CCATCAAGCC	CTGTAAGTCT	GTGTGCGAGC	GGGCCCGGCA	GGGCTGTGAG	420
GTCGTGCTCG (	GGTAGTTCGG	GACATTCAGA	CACACGCTCG	CCCGGGCCGT	CCCGACACTC	
						400
CCCATACTCA '						480
GGGTATGAGT A	AGTTCATGGC	GGTGAGCACC	GGCCTCTTGG	ACCGGACGCT	CCTCGACGGT	
		a	a) aaaa) maa	mma omogogoa	OCCA COMCAM	540
GTGTACGACA						540
CACATGCTGT	CCCCGCACAC	GTAGAGAGGG	CICCGGTAGC	AATGACGCCT	GCCICGACIA	
TTTCCTATGG .	<b>እ</b> ጥጥር ጥ እ ር ጥ እ እ	CCCA A A CTCT	ACACCCCCA A	CCACTGAACG	ርጥርጥል አልጥርጥ	600
AAAGGATACC						• • • • • • • • • • • • • • • • • • • •
MANGGATACC	IMMONICHII	0001110	101000011	00101101100		
AAGCCTATTA	GAGCTACACA	GAAGACCTAT	TTCCGGAACA	ATTACAACTA	TGTCATTCGG	660
TTCGGATAAT						
GCTAAAGTTA						720
CGATTTCAAT	TTCTCTATTT	CTGATTCACG	GTACTACACT	GACGTCATCA	CCTCCACTTC	
						200
					CTATACCAGC	780
CTCTAAGATT	TCAGGAGAGA	CCATTTGTAA	GGTGCCCTG1	GACAGTTGGA	GATATGGTCG	
mamacamaca	mamaaaamaa		አአሙሮአድሮአቶ	ባ <b>አጥአጥ</b> ፖአጥፖአባ	GGGCTATGAA	840
TCTGGCTGCC	ACACCCCTCC	ACTTAATGTT	MAIGAGGAAT MAIGAGGAAT	TATALCATORS	CCCGATACTT	0.10
AGACCGACGG	<b>DEMODED AUM</b>	TGWALIACHE	TIMOTOGIA	' IVIVOIVGIL	. CCCGMINCII	

1

GATGAGGAAC CTACTCCTTG		 			900
CGACTCGGTA GCTGAGCCAT		 			960
AGTGATTCTA					1020
TCACTAAGAT		 			
CGGCAAGCAC GCCGTTCGTG		 			1080
ACTTACTTGC TGAATGAACG		 			1140
	AAAATCATGT TTTTAGTACA	 			1200
	TCTCAACCCC AGAGTTGGGG	 			1260
	TCACTAATCA AGTGATTAGT	 	*		1320
	AGAGCCTCTT TCTCGGAGAA	 			1380
	AATATTGGAT TTATAACCTA	 			1440
	TACTCTGCCG ATGAGACGGC				1500
	TTAGAAAGTT AATCTTTCAA				1560
	GCAAAGCAAT CGTTTCGTTA			ACACCCAAGA TGTGGGTTCT	1620
				AGAACATTTT TCTTGTAAAA	1680
				TAGCATTCTT ATCGTAAGAA	1740
				GAAATGAATT CTTTACTTAA	1800
				AAATTAAATTA T	1860
	AAAGTCAAAA TTTCAGTTTI				